

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/756,095

DATE: 10/24/2001

TIME: 15:56:02

Input Set : A:\09756095seqlisting.txt

Output Set: N:\CRF3\10242001\I756095.raw

3 <110> APPLICANT: Mitchell, Lloyd G.  
4 Garcia-Blanco, Mariano A.  
6 <120> TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR USE IN  
7 SPLICEOSOME MEDIATED RNA TRANS-SPLICING  
10 <130> FILE REFERENCE: A31304-B-A 072874.0134  
12 <140> CURRENT APPLICATION NUMBER: 09/756,095  
13 <141> CURRENT FILING DATE: 2001-01-08  
15 <150> PRIOR APPLICATION NUMBER: 09/158,863  
16 <151> PRIOR FILING DATE: 1998-09-23  
18 <150> PRIOR APPLICATION NUMBER: 09/133,717  
19 <151> PRIOR FILING DATE: 1998-08-13  
21 <150> PRIOR APPLICATION NUMBER: 09/087,233  
22 <151> PRIOR FILING DATE: 1998-05-28  
24 <150> PRIOR APPLICATION NUMBER: 08/766,354  
25 <151> PRIOR FILING DATE: 1996-12-13  
27 <150> PRIOR APPLICATION NUMBER: 60/008,317  
28 <151> PRIOR FILING DATE: 1995-12-07  
30 <160> NUMBER OF SEQ ID NOS: 105  
32 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
34 <210> SEQ ID NO: 1  
35 <211> LENGTH: 132  
36 <212> TYPE: DNA  
37 <213> ORGANISM: Homo sapiens  
39 <400> SEQUENCE: 1  
40 caggggacgc accaaggatg gagatgttcc agggcgctga tgatgttggt gattcttctt 60  
41 aaatcttttg tgatggaaaa cttttcttcg taccacggga ctaaacctgg ttatgtagat 120  
42 tccattcaaa aa 132  
44 <210> SEQ ID NO: 2  
45 <211> LENGTH: 29  
46 <212> TYPE: DNA  
47 <213> ORGANISM: Corynebacterium diptheriae  
49 <400> SEQUENCE: 2  
50 ggcgctgcag ggcgctgatg atgttggtg 29  
52 <210> SEQ ID NO: 3  
53 <211> LENGTH: 36  
54 <212> TYPE: DNA  
55 <213> ORGANISM: Corynebacterium diptheriae  
57 <400> SEQUENCE: 3  
58 ggcgaagctt ggatecgaca cgatttcctg cacagg 36  
60 <210> SEQ ID NO: 4  
61 <211> LENGTH: 68  
62 <212> TYPE: DNA  
63 <213> ORGANISM: Artificial Sequence  
65 <220> FEATURE:  
66 <223> OTHER INFORMATION: Oligonucleotide  
68 <400> SEQUENCE: 4  
69 aattctctag atgcttcacc cgggctgac tcgagtacta actggtacct cttctttttt 60

ENTERED

ENTERED

P-5

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70 ttcctgca 68
72 <210> SEQ ID NO: 5
73 <211> LENGTH: 60
74 <212> TYPE: DNA
75 <213> ORGANISM: Artificial Sequence
77 <220> FEATURE:
78 <223> OTHER INFORMATION: Oligonucleotide
80 <400> SEQUENCE: 5
81 ggaaaaaaa gaagaggtag cagttagtag tcgagtcagg cccgggtgaa gcatctagag 60
84 <210> SEQ ID NO: 6
85 <211> LENGTH: 24
86 <212> TYPE: DNA
87 <213> ORGANISM: Artificial Sequence
89 <220> FEATURE:
90 <223> OTHER INFORMATION: Oligonucleotide primer
92 <400> SEQUENCE: 6
93 tcgagcaacg ttataataat gtgc 24
95 <210> SEQ ID NO: 7
96 <211> LENGTH: 24
97 <212> TYPE: DNA
98 <213> ORGANISM: Artificial Sequence
100 <220> FEATURE:
101 <223> OTHER INFORMATION: Oligonucleotide primer
103 <400> SEQUENCE: 7
104 tcgagaacat tattataacg ttgc 24
106 <210> SEQ ID NO: 8
107 <211> LENGTH: 35
108 <212> TYPE: DNA
109 <213> ORGANISM: Artificial Sequence
111 <220> FEATURE:
112 <223> OTHER INFORMATION: Oligonucleotide primer
114 <400> SEQUENCE: 8
115 aattctctag atcaggcccc ggtgaagcac tcgag 35
117 <210> SEQ ID NO: 9
118 <211> LENGTH: 25
119 <212> TYPE: DNA
120 <213> ORGANISM: Artificial Sequence
122 <220> FEATURE:
123 <223> OTHER INFORMATION: Oligonucleotide primer
125 <400> SEQUENCE: 9
126 tgcttcaccc ggcctgac tagag 25
128 <210> SEQ ID NO: 10
129 <211> LENGTH: 18
130 <212> TYPE: DNA
131 <213> ORGANISM: Homo sapien
133 <400> SEQUENCE: 10
134 tgcttcaccc ggcctga 18
136 <210> SEQ ID NO: 11
137 <211> LENGTH: 16

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138 <212> TYPE: DNA
139 <213> ORGANISM: Homo sapien
141 <400> SEQUENCE: 11
142 ctcttctttt ttttcc 16
144 <210> SEQ ID NO: 12
145 <211> LENGTH: 18
146 <212> TYPE: DNA
147 <213> ORGANISM: Homo sapien
149 <400> SEQUENCE: 12
150 caacgttata ataatgtt 18
152 <210> SEQ ID NO: 13
153 <211> LENGTH: 16
154 <212> TYPE: DNA
155 <213> ORGANISM: Homo sapien
157 <400> SEQUENCE: 13
158 ctgtgattaa tagcgg 16
160 <210> SEQ ID NO: 14
161 <211> LENGTH: 16
162 <212> TYPE: DNA
163 <213> ORGANISM: Homo sapien
165 <400> SEQUENCE: 14
166 cctggacgcg gaagtt 16
168 <210> SEQ ID NO: 15
169 <211> LENGTH: 51
170 <212> TYPE: DNA
171 <213> ORGANISM: Homo sapien
173 <400> SEQUENCE: 15
174 ctgggacaag gacactgctt caccgcgtta gtagaccaca gccctgaagc c 51
176 <210> SEQ ID NO: 16
177 <211> LENGTH: 17
178 <212> TYPE: DNA
179 <213> ORGANISM: Homo sapien
181 <400> SEQUENCE: 16
182 cttctgtttt ttttctc 17
184 <210> SEQ ID NO: 17
185 <211> LENGTH: 16
186 <212> TYPE: DNA
187 <213> ORGANISM: Homo sapien
189 <400> SEQUENCE: 17
190 cttctgtatt attctc 16
192 <210> SEQ ID NO: 18
193 <211> LENGTH: 16
194 <212> TYPE: DNA
195 <213> ORGANISM: Homo sapien
197 <400> SEQUENCE: 18
198 gttctgtcct tgtctc 16
200 <210> SEQ ID NO: 19
201 <211> LENGTH: 29
202 <212> TYPE: DNA

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203 <213> ORGANISM: Corynebacterium diptheriae
205 <400> SEQUENCE: 19
206 ggcgctgcag ggcgctgatg atgttggtg          29
208 <210> SEQ ID NO: 20
209 <211> LENGTH: 36
210 <212> TYPE: DNA
211 <213> ORGANISM: Corynebacterium diptheriae
213 <400> SEQUENCE: 20
214 ggcgaagctt ggatccgaca cgatttcctg cacagg          36
216 <210> SEQ ID NO: 21
217 <211> LENGTH: 21
218 <212> TYPE: DNA
219 <213> ORGANISM: Corynebacterium diptheriae
221 <400> SEQUENCE: 21
222 catcgtcata atttccttgt g          21
224 <210> SEQ ID NO: 22
225 <211> LENGTH: 20
226 <212> TYPE: DNA
227 <213> ORGANISM: Corynebacterium diptheriae
229 <400> SEQUENCE: 22
230 atggaatcta cataaccagg          20
232 <210> SEQ ID NO: 23
233 <211> LENGTH: 20
234 <212> TYPE: DNA
235 <213> ORGANISM: Corynebacterium diptheriae
237 <400> SEQUENCE: 23
238 gaaggctgag cactacacgc          20
240 <210> SEQ ID NO: 24
241 <211> LENGTH: 20
242 <212> TYPE: DNA
243 <213> ORGANISM: Homo sapien
245 <400> SEQUENCE: 24
246 cggcaccgtg gccgaagtgg          20
248 <210> SEQ ID NO: 25
249 <211> LENGTH: 30
250 <212> TYPE: DNA
251 <213> ORGANISM: Homo sapien
253 <400> SEQUENCE: 25
254 accggaattc atgaagccag gtacaccagg          30
256 <210> SEQ ID NO: 26
257 <211> LENGTH: 20
258 <212> TYPE: DNA
259 <213> ORGANISM: Homo sapien
261 <400> SEQUENCE: 26
262 gggcaagggtg aacgtggatg          20
264 <210> SEQ ID NO: 27
265 <211> LENGTH: 19
266 <212> TYPE: DNA
267 <213> ORGANISM: Homo sapien

```

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269 <400> SEQUENCE: 27
270 atcaggagtg gacagatcc 19
272 <210> SEQ ID NO: 28
273 <211> LENGTH: 39
274 <212> TYPE: DNA
275 <213> ORGANISM: Artificial Sequence
277 <220> FEATURE:
278 <223> OTHER INFORMATION: Oligonucleotide primer complimentary to the
279 Escherichia coli lacZ gene
281 <400> SEQUENCE: 28
282 gcatgaattc ggtacccatgg ggggggttctc atcatcatc 39
284 <210> SEQ ID NO: 29
285 <211> LENGTH: 36
286 <212> TYPE: DNA
287 <213> ORGANISM: Artificial Sequence
289 <220> FEATURE:
290 <223> OTHER INFORMATION: Oligonucleotide primer complimentary to the
291 Escherichia coli lacZ gene
293 <400> SEQUENCE: 29
294 ctgaggatcc tcttacctgt aaacgcccac actgac 36
296 <210> SEQ ID NO: 30
297 <211> LENGTH: 38
298 <212> TYPE: DNA
299 <213> ORGANISM: Artificial Sequence
301 <220> FEATURE:
302 <223> OTHER INFORMATION: Oligonucleotide primers complimentary to the
303 Escherichia coli lacZ gene
305 <400> SEQUENCE: 30
306 gcatggtaac cctgcagggc ggcttcgtct gggactgg 38
308 <210> SEQ ID NO: 31
309 <211> LENGTH: 38
310 <212> TYPE: DNA
311 <213> ORGANISM: Artificial Sequence
313 <220> FEATURE:
314 <223> OTHER INFORMATION: Oligonucleotide primers complimentary to the
315 Escherichia coli lacZ gene
317 <400> SEQUENCE: 31
318 ctgaaagctt gttacttat ttttttgac accagacc 38
320 <210> SEQ ID NO: 32
321 <211> LENGTH: 47
322 <212> TYPE: DNA
323 <213> ORGANISM: Artificial Sequence
325 <220> FEATURE:
326 <223> OTHER INFORMATION: Oligonucleotide primers complimentary to the
327 Escherichia coli lacZ gene
329 <400> SEQUENCE: 32
330 gcatggtaac cctgcagggc ggcttcgtct aataatggga ctgggtg 47
332 <210> SEQ ID NO: 33
333 <211> LENGTH: 37

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## VERIFICATION SUMMARY

DATE: 10/24/2001

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TIME: 15:56:03

Input Set : A:\09756095seqlisting.txt

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L:547 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54  
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54  
L:565 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55  
L:566 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55  
L:583 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56  
L:584 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56  
L:828 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:75  
L:950 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:85